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SEQUENCE LISTING

<110> Japan EnviroChemicals, Ltd.

<120> A protein binding to plasticizers

<130> 09622

<150> JP 2003-110877

<151> 2003-4-15

<160> 27

<170> PatentIn version 3.1

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<221> CDS

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Ser Leu Lys Leu Ser Cys Ala Ala Ser Gly Phe Thr Phe Gly Ser Tyr	
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Gly Met Ser Trp Val Arg Gln Thr Ala Asp Lys Arg Leu Glu Trp Val	
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gca acc att tat agt ggt ggt ttt tac acc tac tat cca gac agt gtg	192
Ala Thr Ile Tyr Ser Gly Gly Phe Tyr Thr Tyr Pro Asp Ser Val	
50 55 60	
agg gga cga ttc acc atc tcc aga gac aat gtc aag gaa atc gtg tat	240
Arg Gly Arg Phe Thr Ile Ser Arg Asp Asn Val Lys Glu Ile Val Tyr	
65 70 75 80	
ctg caa atg agc agt ctg aag tct gag gac aca gcc atg tat tac tgt	288
Leu Gln Met Ser Ser Leu Lys Ser Glu Asp Thr Ala Met Tyr Tyr Cys	
85 90 95	
gca aga cgg acg gta gta tct acg gac tat act ttg gac tac tgg ggt	336
Ala Arg Arg Thr Val Val Ser Thr Asp Tyr Thr Leu Asp Tyr Trp Gly	
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gac aga gtc acc atc agt tgc cgg cca agt cag gac atc agc aat ttt 96
Asp Arg Val Thr Ile Ser Cys Arg Pro Ser Gln Asp Ile Ser Asn Phe
20 25 30
tta aac tgg ttt cag cag aaa cca gat gga act gtt gaa gtc ctg atc 144
Leu Asn Trp Phe Gln Gln Lys Pro Asp Gly Thr Val Glu Val Leu Ile
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tgc tac aca tta aga atg cac tta gga gtc cca tca acg ttc agt ggc 192
Cys Tyr Thr Leu Arg Met His Leu Gly Val Pro Ser Thr Phe Ser Gly
50 55 60
tgt gtg tct gga aca tat tat act ctc acc agt agc aac ctg gaa caa 240
Cys Val Ser Gly Thr Tyr Tyr Thr Leu Thr Ser Ser Asn Leu Glu Gln
65 70 75 80
gaa gat ata gac act tcc ttt gcc att agg att ata cgc gtg ctc acg 288
Glu Asp Ile Asp Thr Ser Phe Ala Ile Arg Ile Ile Arg Val Leu Thr
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Val Gly Ala Gly Thr Thr Leu Glu Leu Lys
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Cys Tyr Thr Leu Arg Met His Leu Gly Val Pro Ser Thr Phe Ser Gly
50 55 60
Cys Val Ser Gly Thr Tyr Tyr Thr Leu Thr Ser Ser Asn Leu Glu Gln
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Lys Gly

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 Ser Leu Ser Leu Thr Cys Ser Val Thr Gly Tyr Ser Ile Thr Ser Gly
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tat tac tgg aat tgg atc cgg caa ttt oca gga aac aaa ctg gat tgg 144
 Tyr Tyr Trp Asn Trp Ile Arg Gln Phe Pro Gly Asn Lys Leu Asp Trp
 35 40 45

atg ggc cat ata agt tac gac ggt aac aat aac tac aac cca tct ctc 192
 Met Gly His Ile Ser Tyr Asp Gly Asn Asn Asn Tyr Asn Pro Ser Leu
 50 55 60

aaa aat cga atc tcc atc act cgt gac aca tct aag aac cag ttt ttc 240
 Lys Asn Arg Ile Ser Ile Thr Arg Asp Thr Ser Lys Asn Gln Phe Phe
 65 70 75 80

ctg aag ttg aat tct gtg act act gag gac aca gat aca tat tac tgt 288
 Leu Lys Leu Asn Ser Val Thr Thr Glu Asp Thr Asp Thr Tyr Tyr Cys
 85 90 95

tct atg atc ctc tat ggt atg gac tac tgg ggt cag gga acc tca gtc 336
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 35 40 45
 Met Gly His Ile Ser Tyr Asp Gly Asn Asn Asn Tyr Asn Pro Ser Leu
 50 55 60
 Lys Asn Arg Ile Ser Ile Thr Arg Asp Thr Ser Lys Asn Gln Phe Phe
 65 70 75 80
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 gaa cgg gtc acc atg acc tgc act gcc agc tca agt gta agt tcc agt 96
 Glu Arg Val Thr Met Thr Cys Thr Ala Ser Ser Ser Val Ser Ser Ser
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 tac ttg cac tgg tac cag cag aag cca gga tcc tcc ccc aaa ctc tgc 144
 Tyr Leu His Trp Tyr Gln Gln Lys Pro Gly Ser Ser Pro Lys Leu Cys
 35 40 45
 att tat agc aca tcc aac ctg gct tct gga gtc cca act cgc ttc agt 192
 Ile Tyr Ser Thr Ser Asn Leu Ala Ser Gly Val Pro Thr Arg Phe Ser
 50 55 60
 ggc agt ggg tct ggg acc tct tac tct ctc aca ata agc agc atg gag 240
 Gly Ser Gly Ser Gly Thr Ser Tyr Ser Leu Thr Ile Ser Ser Met Glu
 65 70 75 80
 gct gaa gat gct gcc act tat tac tgc cac cag tat cat cgt tcc cca 288
 Ala Glu Asp Ala Ala Thr Tyr Tyr Cys His Gln Tyr His Arg Ser Pro
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 35 40 45
 Ile Tyr Ser Thr Ser Asn Leu Ala Ser Gly Val Pro Thr Arg Phe Ser
 50 55 60

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Ala	Glu	Asp	Ala	Ala	Thr	Tyr	Tyr	Cys	His	Gln	Tyr	His	Arg	Ser	Pro
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